

58C

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/518,181
Source: PCT
Date Processed by STIC: 01/05/2006

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PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/518,181

DATE: 01/05/2006
TIME: 11:04:34

Input Set : A:\07947.ST25.txt
Output Set: N:\CRF4\01052006\J518181.raw

3 <110> APPLICANT: The Regents of the University of California
 5 <120> TITLE OF INVENTION: Methods for Inhibiting Angiogenesis, Cell Migration, Cell
 6 Adhesion, and Cell Survival
 8 <130> FILE REFERENCE: UCSD-07947
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/518,181
 C--> 10 <141> CURRENT FILING DATE: 2004-12-15
 10 <160> NUMBER OF SEQ ID NOS: 116
 12 <170> SOFTWARE: PatentIn version 3.2
 14 <210> SEQ ID NO: 1
 15 <211> LENGTH: 68
 16 <212> TYPE: DNA
 17 <213> ORGANISM: Artificial Sequence
 19 <220> FEATURE:
 20 <223> OTHER INFORMATION: Synthetic
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 25 cattgtat 68
 28 <210> SEQ ID NO: 2
 29 <211> LENGTH: 102
 30 <212> TYPE: PRT
 31 <213> ORGANISM: Artificial Sequence
 33 <220> FEATURE:
 34 <223> OTHER INFORMATION: Synthetic
 36 <400> SEQUENCE: 2
 38 Asp Thr Val Lys Val Met Val Val Met Leu Ala Ile Cys Phe Leu Ala
 39 1 5 10 15
 42 Arg Ser Asp Gly Lys Pro Val Lys Lys Arg Ser Val Ser Glu Ile Gln
 43 20 25 30
 46 Leu Met His Asn Leu Gly Lys His Leu Ser Ser Leu Glu Arg Val Glu
 47 35 40 45
 50 Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn Phe Val Val Leu Gly
 51 50 55 60
 54 Ala Ser Ile Val His Arg Asp Gly Gly Ser Gln Arg Pro Pro Lys Lys
 55 65 70 75 80
 58 Glu Asp Asn Val Leu Val Glu Ser His Gln Lys Ser Leu Gly Glu Ala
 59 85 90 95
 62 Asp Lys Ala Ala Val Gly
 63 100
 66 <210> SEQ ID NO: 3
 67 <211> LENGTH: 306
 68 <212> TYPE: DNA
 69 <213> ORGANISM: Artificial Sequence
 71 <220> FEATURE:

(pg-6)

RAW SEQUENCE LISTING

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Output Set: N:\CRF4\01052006\J518181.raw

72 <223> OTHER INFORMATION: Synthetic
 74 <400> SEQUENCE: 3
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 77 aaggctgtta agaagagatc tggagatggaa atacagctta tgcataaccc gggcaaacac 120
 79 ctgagctctc tggagagatg ggaatggctg cgaaagaagc tgcaggatgt gcacaacttt 180
 81 gttgttctcg gagcttctat agttcacaga gatggtggtt cccagagacc cccaaaaaaag 240
 83 gaagacaatg tccttagttga gagccatcaa aaaagtctcg gagaagcaga taaagctgct 300
 85 gtgggg 306

88 <210> SEQ ID NO: 4
 89 <211> LENGTH: 67
 90 <212> TYPE: PRT
 91 <213> ORGANISM: Artificial Sequence
 93 <220> FEATURE:
 94 <223> OTHER INFORMATION: Synthetic
 96 <400> SEQUENCE: 4
 98 Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn
 99 1 5 10 15
 102 Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln
 103 20 25 30
 106 Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys
 107 35 40 45
 110 Ser Leu Gly Glu Ala Asp Lys Ala Asp Val Asn Val Leu Thr Lys Ala
 111 50 55 60
 114 Lys Ser Gln
 115 65
 118 <210> SEQ ID NO: 5
 119 <211> LENGTH: 290
 120 <212> TYPE: DNA
 121 <213> ORGANISM: Artificial Sequence
 123 <220> FEATURE:
 124 <223> OTHER INFORMATION: Synthetic
 126 <400> SEQUENCE: 5
 127 aattctgtaa gggatccaag aagagatcg tttctgagat ccaattgtat cataacttgg 60
 129 gtaagcaattt gaaactctatg gaaagatgg aatgggttgcgtt aaagaagatcg caggacgttc 120
 131 acaacttcgt tgctttggaa gctccattgg ctccaaagaga cgctgggttcaaagaccaa 180
 133 gaaagaagggaa agacaacgtt ttgggttgcgtt ctcacggaaa gtctttgggtt gaagctgaca 240
 135 aggctgacgt taacgtgtta actaaggctt aatcgcaata aagatcttga 290
 138 <210> SEQ ID NO: 6
 139 <211> LENGTH: 120
 140 <212> TYPE: DNA
 141 <213> ORGANISM: Homo sapiens
 143 <400> SEQUENCE: 6
 144 aggagctcca agggcaacaa aattgtgcac atcctgcagc ttcttacgca gccattctac 60
 146 tctctccatc gagttcagat gttttccatc gttatgcata agctgtattt cactcacaga 120
 149 <210> SEQ ID NO: 7
 150 <211> LENGTH: 116
 151 <212> TYPE: DNA
 152 <213> ORGANISM: Homo sapiens
 154 <400> SEQUENCE: 7

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155 gagtcatcat ccattttcca ttttcctct ccatccacaa gcataatactc aatattcgt 60
157 cttttttttt ttttttttg agacagggtc tcactctgtc gcccaggctg gagtac 116
160 <210> SEQ ID NO: 8
161 <211> LENGTH: 115
162 <212> TYPE: PRT
163 <213> ORGANISM: Homo sapiens
165 <400> SEQUENCE: 8
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168 1 5 10 15
171 Ile Cys Phe Leu Thr Lys Ser Asp Gly Lys Ser Val Lys Lys Arg Ser 30
172 20 25 30
175 Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn Ser 45
176 35 40 45
179 Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn 60
180 50 55 60
183 Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln 80
184 65 70 75 80
187 Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys 95
188 85 90 95
191 Ser Leu Gly Glu Ala Asp Lys Ala Asp Val Asn Val Leu Thr Lys Ala 110
192 100 105
195 Lys Ser Gln
196 115
199 <210> SEQ ID NO: 9
200 <211> LENGTH: 772
201 <212> TYPE: DNA
202 <213> ORGANISM: Homo sapiens
204 <400> SEQUENCE: 9
205 tgtctttagt ttactcagca tcagctacta acataacctga acgaagatct tggtaaga 60
207 cattgtatgt gaagatgata cctgcaaaag acatggctaa agttatgatt gtcattgtgg 120
209 caatttgttt tcttacaaaa tcggatggga aatctgttaa gaagagatct gtgagtgaaa 180
211 tacagcttat gcataacctg ggaaaaacatc tgaactcgtat ggagagagta gaatggctgc 240
213 gtaagaagct gcaggatgtg cacaattttg ttgccttgg agctcctcta gctccagag 300
215 atgctgggtc ccagaggccc cgaaaaaagg aagacaatgt cttgggttag agccatgaaa 360
217 aaagtcttgg agaggcagac aaagctgtat tgaatgtatt aactaaagct aaatcccagt 420
219 gaaaatgaaa acagatattt ttagatgtt cttttttttt gttttttttt aatgtatgtt 480
221 ctgctaattc aaagcttatc taagatttttcc aagtccat atttctgata taacaaacta 540
223 catgttatcc atcactatgcc atgataactt cttttttttt tgattttttt gttttttttt 600
225 ttattcattt gagttatccc aattatcttt tctattgtttt attctttttt aatgtatgtt 660
227 ttgcataatt tataaaagaa taaaatttgc cttttttttt tttttttttt aatgtatgtt 720
229 aaaacaaaaaa tgtaatgtatc ataagtctaa ataaatgtatc tttttttttt tttttttttt 772
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233 <211> LENGTH: 27
234 <212> TYPE: DNA
235 <213> ORGANISM: Unknown
237 <220> FEATURE:
238 <223> OTHER INFORMATION: Synthetic
240 <400> SEQUENCE: 10
241 qatcctcatt actqqqattt aqcttta 27

RAW SEQUENCE LISTING

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Output Set: N:\CRF4\01052006\J518181.raw

244 <210> SEQ ID NO: 11
 245 <211> LENGTH: 42
 246 <212> TYPE: DNA
 247 <213> ORGANISM: Unknown
 249 <220> FEATURE:
 250 <223> OTHER INFORMATION: Synthetic
 252 <400> SEQUENCE: 11
 253 taaaaaatcc ctaggcagg cagacaaggc cgatgtaat gt 42
 256 <210> SEQ ID NO: 12
 257 <211> LENGTH: 263
 258 <212> TYPE: DNA
 259 <213> ORGANISM: Artificial Sequence
 261 <220> FEATURE:
 262 <223> OTHER INFORMATION: Synthetic
 264 <400> SEQUENCE: 12
 265 tatgtctgtc tccgagattc agttaatgca taaccctggc aaacatttga actccatgg 60
 267 gcgtgtagaa tggctcgta agaagttgca ggatgtgcac aattttgttg ccttaggtgc 120
 269 cccattggct cctcgtgatg ctggttccca aagaccacgt aaaaaggaag acaatgtctt 180
 271 agttgagagc catgaaaaat cccttaggcga ggcagacaag gccgatgtga atgtattaac 240
 273 taaagctaaa tcccagtaat gag 263
 276 <210> SEQ ID NO: 13
 277 <211> LENGTH: 614
 278 <212> TYPE: DNA
 279 <213> ORGANISM: Artificial Sequence
 281 <220> FEATURE:
 282 <223> OTHER INFORMATION: Synthetic
 284 <400> SEQUENCE: 13
 285 tttgagttagaa aataacttca tttatatttca cttatgtatca ttacattttt gttttacatt 60
 287 ttaaggtaga agagaggttt aaaagtgcggaa ttttatttctt ttataaatta tgcaataaca 120
 289 tactttaaaa agaataaaaca atagaaaaga taattttaaat aactcaaatg aataaaaatg 180
 291 gaatcagaat aatcaattaa aattgcaggat atcatggctt gtatggatt acatgttagtt 240
 293 ttttatatca gaaatattgg cacttggaaa tcttaataga gctttgaatt agcagcatgt 300
 295 atttttgcggcc tacactgtctt agagcagaac tctgacaata tctgttttca ttttcactgg 360
 297 gattttagctt tagtttaatac attcacatca gctttgtctt cctctccaatg actttttca 420
 299 tggctctcaa ccaagacatt gtcttccttt tttcggggcc tctggaaacc agcatcttg 480
 301 ggagcttagag gagctccaatggcacaacaaa ttgtgcacat cctgcagatgtt cttacgcac 540
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 305 ctcacagatctt 614
 308 <210> SEQ ID NO: 14
 309 <211> LENGTH: 68
 310 <212> TYPE: DNA
 311 <213> ORGANISM: Unknown
 313 <220> FEATURE:
 314 <223> OTHER INFORMATION: Synthetic
 316 <400> SEQUENCE: 14
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 319 taaagaca 68
 322 <210> SEQ ID NO: 15
 323 <211> LENGTH: 115

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324 <212> TYPE: PRT
325 <213> ORGANISM: Homo sapiens
327 <400> SEQUENCE: 15
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330 1 5 10 15
333 Ile Cys Phe Leu Thr Lys Ser Asp Gly Lys Ser Val Lys Lys Arg Ser
334 20 25 30
337 Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn Ser
338 35 40 45
341 Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn
342 50 55 60
345 Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln
346 65 70 75 80
349 Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys
350 85 90 95
353 Ser Leu Gly Glu Ala Asp Lys Ala Asp Val Asn Val Leu Thr Lys Ala
354 100 105 110
357 Lys Ser Gln
358 115
361 <210> SEQ ID NO: 16
362 <211> LENGTH: 1156
363 <212> TYPE: DNA
364 <213> ORGANISM: Homo sapiens
366 <400> SEQUENCE: 16
367 aagcttctcg taaaaaccaa cccaaattgt tagtattgca ttctgtgtac tatagtttgg 60
369 aatattaaaa atatttaaaa atacctccat tttgcttatac ctttttagtga agatgataacc 120
371 tgcaaaaagac atggctaaag ttatgattgt catgttggca atttggtttc ttacaaaatc 180
373 gcatggaaaa tctgttaagt aagtactgtt ttgccttgga attggatttt taatgttgac 240
375 ttatcattt cgaagtgggg agctaattggg aagtggccct ctctgttct cttcttccca 300
377 ggaagagatc tgcgttgcataatc tgcataaccc gggaaaacat ctgaactcga 360
379 tggagagatc agaatggctg cgtaagaagc tgcaggatgt gcacaatttt gttggcccttg 420
381 gagctcctct agctcccaga gatgctggg cccagaggcc ccggaaaaaaag gaagacaatg 480
383 tcttgggttga gagccatgaa aaaagtcttgg gagaggcaga caaagctgtat gtgaatgtat 540
385 taactaaagc taaatcccg taaaaatgaa aacagatatt gtcagagttc tgctctagac 600
387 agttagggc aacaatacat gctgctaatt caaagctcta ttaagatttc caagtgc当地 660
389 tatttctgtat ataacaaact acatgtatcatc catcaacttgc catgataact gcaattttaa 720
391 ttgatttatttc tgattccact tttatttatttc tgaggatttt taatttatttc ttcttatttt 780
393 tattttttt aaagtatgtt attgcataat ttataaaaaga ataaaattgc acttttaaac 840
395 ctctcttcta ccttaaaaatg taaaacaaaa atgtatgtat cataagtcta aataaaatgaa 900
397 gtatttctca ctcattgcaa gtatattttt ttgggttatca ctgataaccc catgtttaca 960
399 ttgatcatgaa cttaggtgaa caatacacaag tatttttttgcatgtgtt tcacattttgg 1020
401 atattttgaa catcaacgtt tttagtattac caaagtatta ggtttccaaa tcttcacttag 1080
403 ctcataactg ttgtccctttt ggtttcagga aaggaaataa aatgctcagc aaaaaaaagg 1140
405 ggcataaaaag tggacc 1156
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409 <211> LENGTH: 526
410 <212> TYPE: DNA
411 <213> ORGANISM: Homo sapiens
413 <400> SEQUENCE: 17

RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:45; N Pos. 426, 427, 428, 429, 430, 438, 439, 440, 441, 442, 443, 449, 450, 451, 492

Seq#:45; N Pos. 493, 494

Seq#:90; N Pos. 8, 23, 32, 48, 59

Seq#:114; Xaa Pos. 10

VERIFICATION SUMMARY

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1553 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:420
L:1555 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:480
L:4242 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:90 after pos.:0
L:5604 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:114 after pos.:0